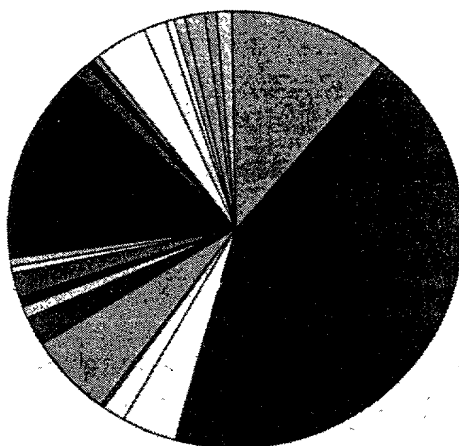


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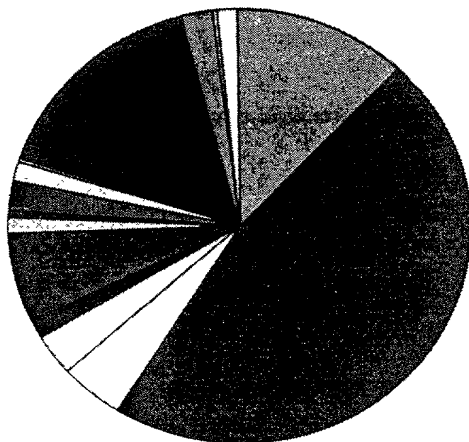
Fig. 1

EG+ cell profile 11/99



- ☒ Housekeeping
- ☒ Unknown
- ☒ Novel
- ☒ Cell cycle
- ☒ Proteinase
- ☒ Cytoskeletal
- ☒ Signal transduction/signaling
- ☒ Protein kinase/bp
- ☒ Protein phosphatase
- ☒ Energy metabolism/biosynthesis
- ☒ Apoptosis
- ☒ Endocrine
- ☒ Transcription factors
- ☒ Other
- ☒ Stress related genes
- ☒ Membrane proteins
- ☒ Protein degradation pathways
- ☒ ECM
- ☒ Viral transposable elements
- ☒ Growth factors/cytokines
- ☒ Protein transport/trafficking
- ☒ Wound healing
- ☒ Cell adhesion/junction
- ☒ Development/Differentiation

ES(-) cell profile 11/99



- ☒ Housekeeping
- ☒ Unknown
- ☒ Novel
- ☒ Cell cycle
- ☒ Proteinases
- ☒ Cytoskeletal
- ☒ Signal transduction/signaling
- ☒ Protein kinases/bp
- ☒ Protein phosphatase
- ☒ Energy metabolism/biosynthesis
- ☒ Apoptosis
- ☒ Endocrine
- ☒ Transcription factor
- ☒ Other
- ☒ Stress related proteins
- ☒ Membrane proteins
- ☒ Protein degradation pathways
- ☒ ECM
- ☒ Viral transposable elements
- ☒ Growth factors/cytokines
- ☒ Protein transport/trafficking
- ☒ Wound healing
- ☒ Cell junction/adhesion
- ☒ Development/Differentiation

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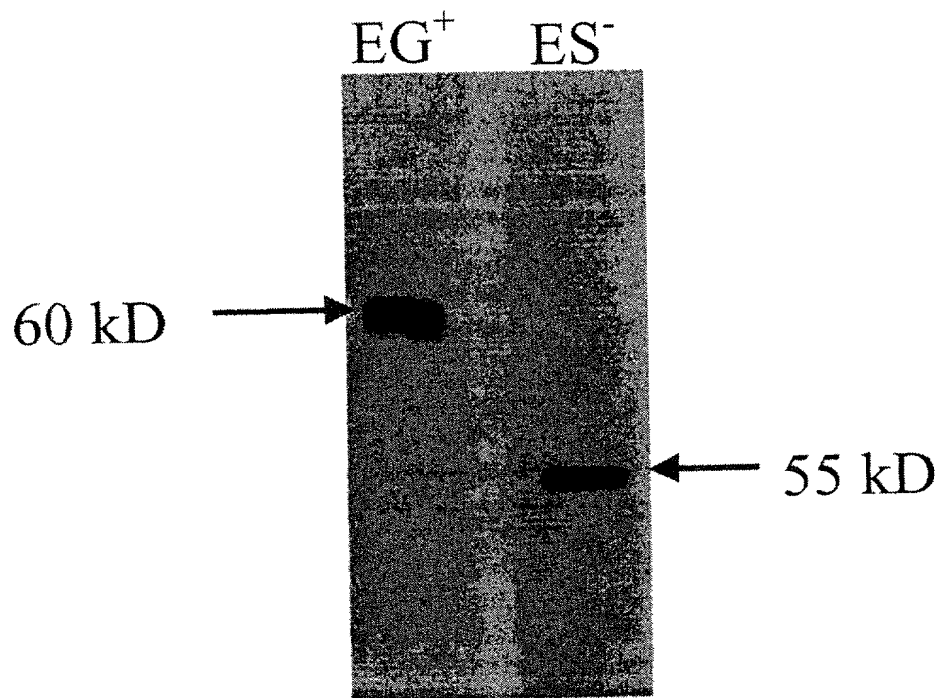


FIG. 2

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P5/T5
IV EG ES

P6/T4
IV EG ES

P2/T9
IV EG ES

P4/T7
IV EG ES

P3/T8
IV EG ES

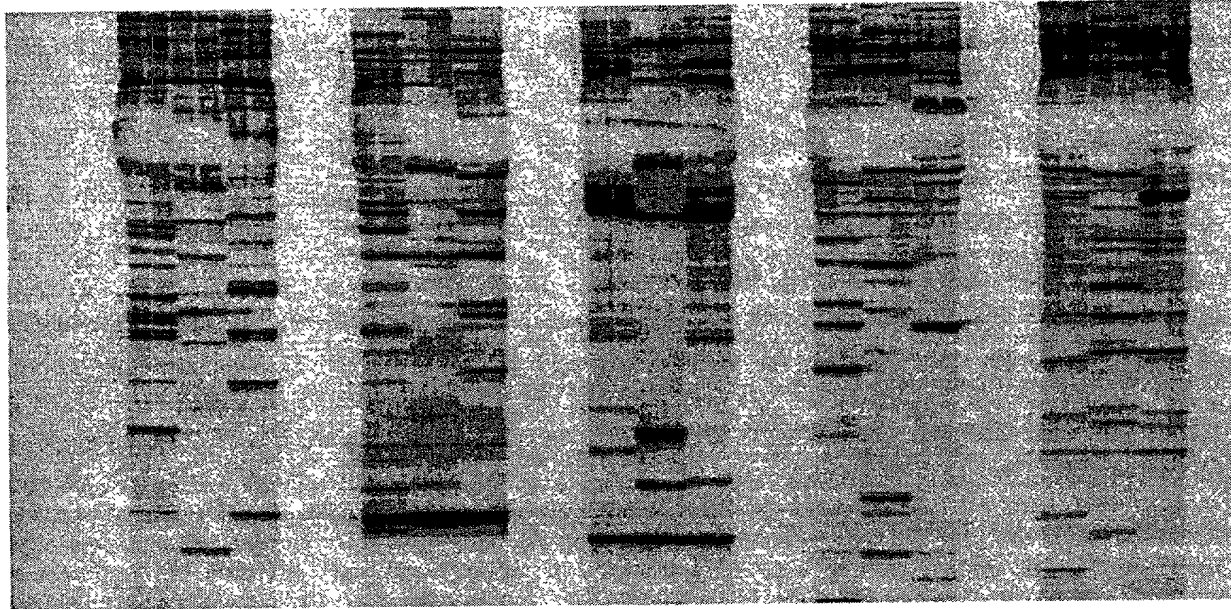


FIG. 3

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1 2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 (DC)-Donor Cell

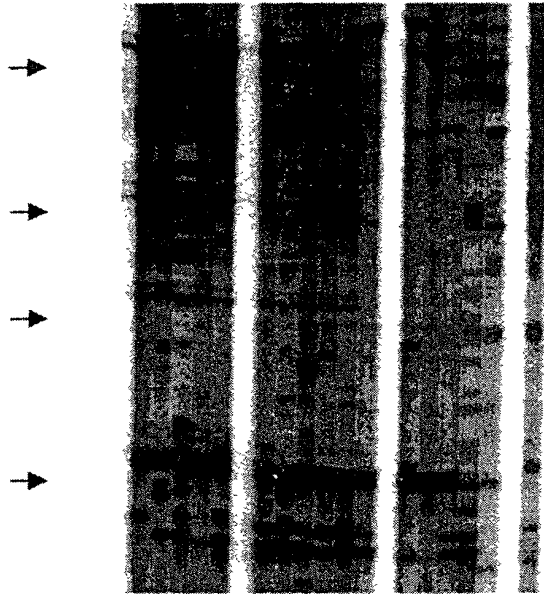
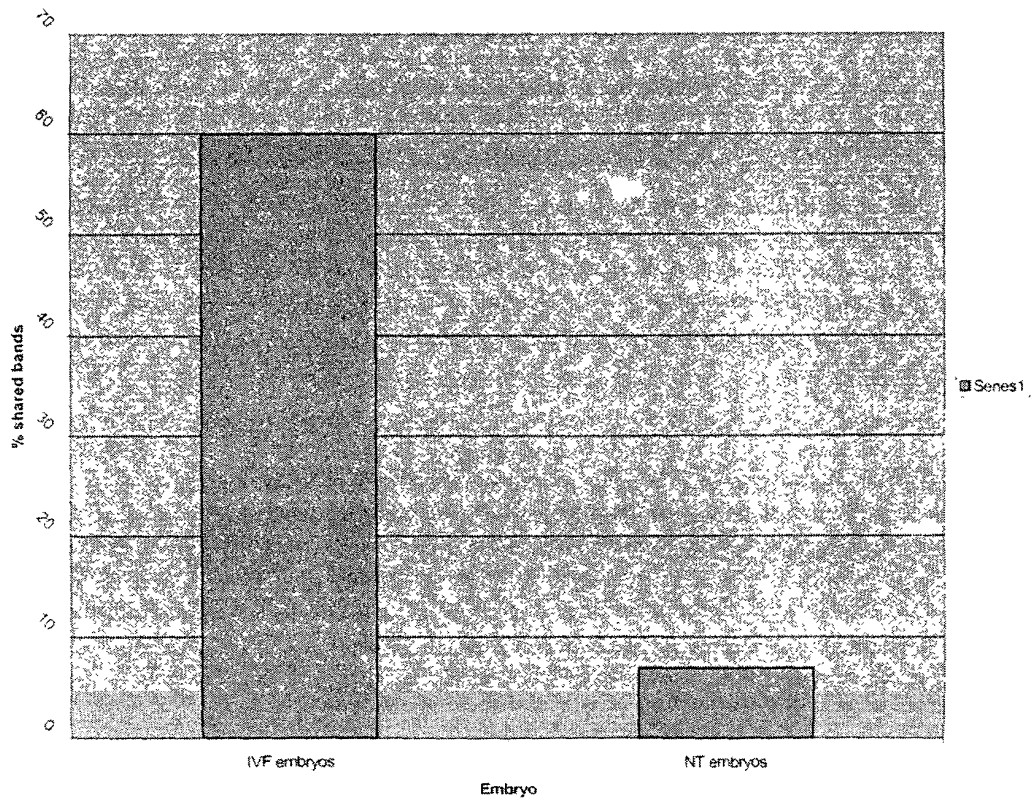


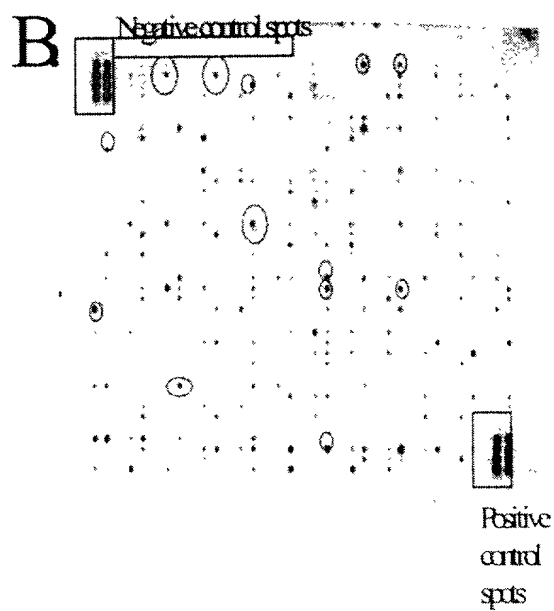
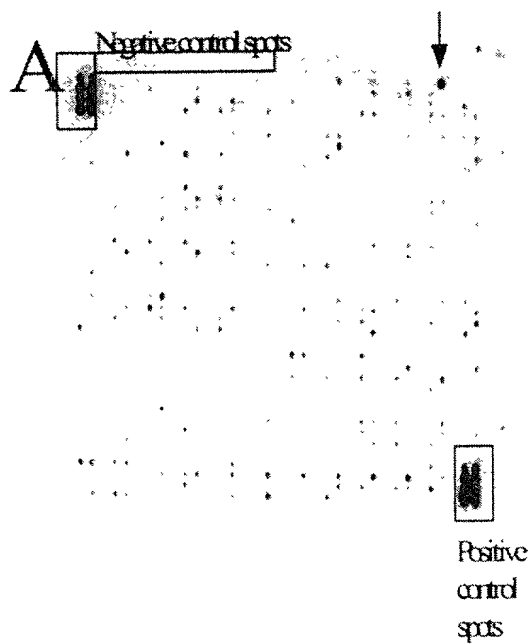
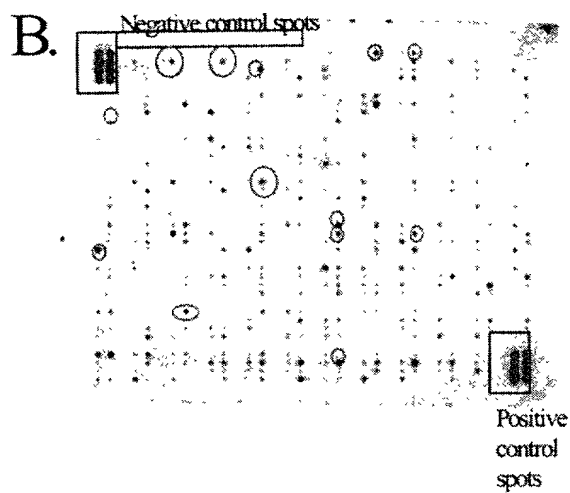
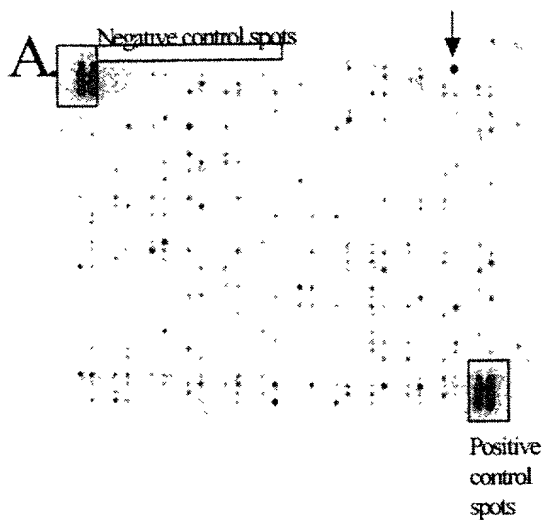
Fig. 4a

Fig. 4b

Comparison of DD Banding Patterns



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09076143.02602

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3110
307 11 831

2008220-CHT92860

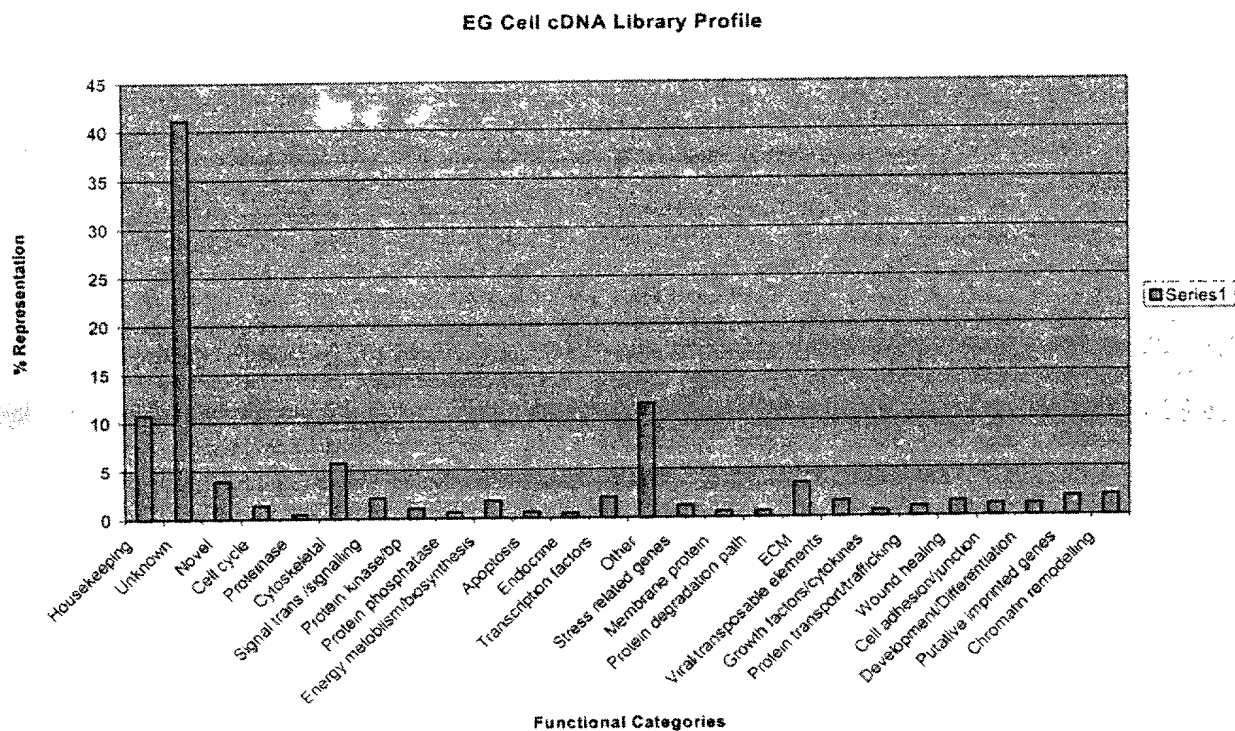


Fig. 6

09076143.022602

3910
196 13 93

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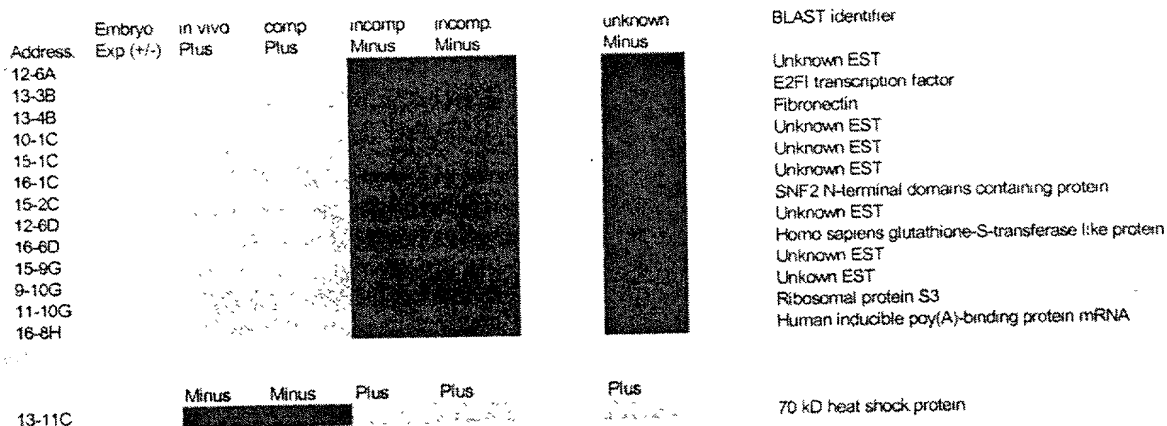


FIG. 7